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4
5 ***Cucumis metuliferus* is resistant to root-knot nematode *Mi1.2* gene (a)virulent**
6 **populations and a promising melon rootstock.**

7
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18
19 ABSTRACT

20 Pot experiments were carried out to characterize the response of two *Cucumis metuliferus*
21 accessions against (a)virulent *Meloidogyne arenaria*, *M. incognita* or *M. javanica*
22 populations, to *Mi1.2* gene and to determine the compatibility and the effect on
23 physicochemical properties of cantaloupe melon. In addition, histopathological studies
24 were conducted. Plants were inoculated in 200 cm³-pots with 1 J2 cm⁻³ of soil containing

25 sterilized sand a week after transplanting and maintained in a growth chamber at 25 °C
26 for 40 days. The susceptible cucumber cv. Dasher II or melon cv. Paloma were included
27 for comparison. The number of egg masses and number of eggs per plant were assessed,
28 and the reproduction index (RI) was calculated as the percentage of eggs produced on the
29 *C. metuliferus* accessions respect those produced on the susceptible cultivars. The
30 compatibility and fruit quality was assessed grafting three scions (two of Charentais type)
31 and one of type Piel de Sapo under commercial greenhouse conditions. The resistance
32 level of both *C. metuliferus* accessions ranged from highly (RI < 1%) to resistant (1% ≤
33 RI ≤ 10%) irrespective of *Meloidogyne* populations. Melon plants grafted onto *C.*
34 *metuliferus* accession BGV11135 grew as selfgrafted plants and did not modify
35 negatively fruit quality traits. Giant cells induced by RKN on *C. metuliferus* were mostly
36 poor developed compared to those on cucumber. Furthermore, necrotic areas surrounding
37 the nematode were observed. *C. metuliferus* accession BGV11135 could be a promising
38 melon rootstock to manage *Meloidogyne spp.* irrespective of its (a)virulent *Mi1.2*
39 condition without melon fruit quality reduction.

40

41 Key words: *Cucumis melo*, grafting, histopathology, horned cucumber, *Meloidogyne*,
42 plant resistance.

43

44 INTRODUCTION

45 Root-knot nematodes (RKN), *Meloidogyne spp.*, are the most limiting plant parasitic
46 nematodes for vegetable production worldwide (Sikora & Fernández, 2005). Nonetheless,
47 the capability to any RKN species to use a given plant species, to reproduce on it, and to
48 affect its productivity differs according to its host status. Regarding cucurbit crops, one
49 of the most widely cultivated groups around the world, zucchini-squash and watermelon

50 are susceptible and poor-host, respectively, but both are tolerant (López-Gómez *et al.*,
51 2015 & 2016), whilst melon and cucumber are susceptible and are severely damaged by
52 RKN (Di Vito *et al.*, 1983; Giné *et al.*, 2014 & 2017). In Spain, crop rotation
53 sequences including solanaceous and cucurbit crops are very common (Ornat *et al.*, 1997;
54 Talavera *et al.*, 2012; Giné *et al.*, 2016), but there are not available commercial resistant
55 cucurbit cultivars or rootstocks. A way to suppress RKN populations and to reduce yield
56 losses of the most susceptible-intolerant cucurbit crops by non-chemical methods,
57 according to the European directive 2009/128/CE, is grafting onto resistant-tolerant
58 rootstocks. Plant resistance is an effective and profitable control method (Sorribas *et al.*,
59 2005) reducing the RKN reproduction rate and the equilibrium density (Talavera *et al.*,
60 2009; Giné & Sorribas, 2017), and thus, the subsequent yield losses for the next crop
61 (Ornat *et al.*, 1997) which are directly related to nematode population densities in the soil
62 at planting (Seinhorst, 1965). Grafting is an effective tool for controlling other soil borne
63 pathogens (Lee *et al.*, 2010). In this sense, cucurbit crops are usually grafted onto
64 *Cucurbita* hybrids, which are resistant to fusarium wilt but susceptible to *Meloidogyne*
65 *spp.* (Thies *et al.*, 2010; López-Gómez *et al.*, 2016; Giné *et al.*, 2017). However,
66 resistance to RKN has been found in wild *Cucumis* spp., including accessions of *Cucumis*
67 *africanus*, *Cucumis anguria*, *C. ficifolius*, *C. metuliferus*, *C. myriocarpus*, *C. postulatus*,
68 *C. subsericeus*, and *C. zeyheri* (Fassuliotis, 1967; Sigüenza *et al.*, 2005; Kokalis-Burelle
69 & Roskopf, 2011; Pofu *et al.*, 2011; Thies *et al.*, 2014; Liu *et al.*, 2015). Moreover, some
70 of these *Cucumis* species are resistant to pathogenic fungi, such as *Fusarium oxysporum*
71 f. sp. *melonis* (Liu *et al.*, 2015) and *Monosporascus cannonballus* (Dias *et al.*, 2001). The
72 inclusion of RKN resistant cucurbit rootstocks in the solanaceous-cucurbitaceous rotation
73 sequence could be helpful to manage virulent nematode populations to *Mil.2* resistance
74 gene on tomato, which have been increased in the last years (Tzortzakakis *et al.*, 2005;

75 Devran & Sögüt, 2010; Verdejo-Lucas *et al.*, 2012). Nonetheless, as far we know, there
76 is no information about the host suitability of *C. metuliferus* accessions to *Mil.2* virulent
77 RKN populations.

78 *Cucumis metuliferus* is a compatible rootstock for melon but can affect fruit
79 quality traits such as total soluble solids content (° Brix) and flesh firmness depending on
80 melon type and agronomic conditions (Guan *et al.*, 2014). Then, the evaluation for quality
81 traits in different scions is convenient when testing for putative rootstocks. The objective
82 of this study was to assess the host suitability of *C. metuliferus* against RKN (a)virulent
83 populations, its compatibility with melon and the effects on fruit quality. In addition,
84 histopathological studies were conducted to identify resistance mechanisms of *C.*
85 *metuliferus* against *M. javanica*.

86

87 MATERIALS AND METHODS

88 *Nematode inoculum*

89 RKN populations belonging to *M. arenaria*, *M. incognita* and *M. javanica* were used in
90 the experiments. The information on RKN species, code, origin and the (a) virulent status
91 against tomato cultivars carrying the *Mil.2* gene is presented in Table 1. The RKN
92 populations were maintained on the susceptible tomato cv. Durinta (Semini Seeds).
93 Second stage juveniles (J2) were used as inoculum. Eggs were extracted from tomato
94 roots by blender maceration in a 5% of commercial bleach (40 g L⁻¹ NaOCl) solution for
95 5 min (Hussey & Barker, 1973). The egg suspension was then passed through a 74 µm
96 aperture sieve to remove root debris, and eggs were collected on a 25 µm sieve and placed
97 on Baermann trays (Whitehead & Hemming, 1965) at 25 °C. Nematodes were collected
98 daily using a 25 µm sieve during 7 days and stored at 9 °C until inoculation. *Meloidogyne*

99 species were identified according to the morphology of the perineal pattern of the females,
100 and by SCAR-PCR markers (Zijlstra *et al.*, 2000).

101 *Plant material*

102 In the experiments conducted to assess the response of *C. metuliferus* against (a)virulent
103 RKN populations, accessions BGV11135 and BGV10762 of *C. metuliferus*, from the
104 COMAV-UPV collection (Valencia, Spain), and the susceptibles cucumber cv. Dasher II
105 (Semini Seeds) or melon cv. Paloma (Fitó) were used. Seeds of *C. metuliferus* were
106 surface disinfested using a 20% bleach commercial solution (40g L⁻¹ NaOCl) during 2
107 min and washed two times in sterilized distilled water to remove bleach. Afterwards,
108 seeds were placed in Petri dishes with a cotton matrix, irrigated and incubated two days
109 at 37 °C. After germination, seeds were transferred to a tray containing sterile vermiculite,
110 covered with 1.5 cm with the same substrat and placed in a growth chamber at 25±2 °C
111 and 16:8 h (light:darkness) photoperiod for a week. After that, seedlings were individually
112 transplanted to 200 cm³ pots containing sterile river sand. Plants were fertilized with a
113 slow release fertilizer (15% N + 9% P₂O₅ + 12% K₂O + 2% MgO₂ + microelements;
114 Osmocote Plus).

115 To assess *C. metuliferus* compatibility with melon, Charentais melon (*Cucumis*
116 *melo* L. var. *cantalupensis* Naudin) cv. Vedrantaís (COMAV) and cv. Paloma (Fitó), and
117 Piel de sapo melon (*Cucumis melo* L. var. *inodorus* Naudin) cv. Finura (Rijk Zwaan) were
118 used.

119 *Response of C. metuliferus accessions to avirulent RKN populations*

120 Two experiments were carried out to evaluate the response of *C. metuliferus* against
121 avirulent RKN populations. In the first experiment, accessions BGV11135 and
122 BGV10762 of *C. metuliferus*, and cucumber cv. Dasher II were inoculated with 1 J2 cm⁻
123 ³ of soil of the *M. incognita* population Agropolis or the *M. javanica* population MJ05.

124 Plants were maintained in a growth chamber at 25 ± 2 °C and 16:8 h (light:darkness)
125 photoperiod for 40 days. Plants were watered as needed during the experiment. Each
126 plant-RKN population combination was replicated 10 times. Soil temperatures were
127 recorded daily at 30-min interval with a PT100 probe (Campbell Scientific Ltd) placed
128 into the pots at 4 cm depth. At the end of the experiment, roots were carefully washed,
129 weighted and immersed in a 0.01% solution of erioglaucina to stain egg masses in blue
130 (Omwega *et al.*, 1988) previous to count them. RKN eggs were extracted from roots by
131 maceration in a 10% bleach commercial solution (40g L^{-1} NaOCl) (Hussey & Barker,
132 1973). The number of eggs was counted and the reproduction index (RI) was calculated
133 as the percentage of the number of eggs per plant in the experimental accession with
134 respect to that on the susceptible cucumber cv. Dasher II. After that, the response of the
135 accessions was categorized according to the RI as, highly resistant ($\text{RI} < 1\%$), resistant
136 ($1\% \leq \text{RI} < 10\%$), moderately resistant ($10\% \leq \text{RI} < 25\%$), slightly resistant ($25\% \leq \text{RI} <$
137 50%) or susceptible ($\text{RI} \geq 50\%$) (Hadisoeganda & Sasser, 1982).

138 In the second experiment, the response of the *C. metuliferus* accession BGV11135
139 and the susceptible standard cucumber cv. Dasher II was assessed against one population
140 of *M. arenaria* (MA68), two populations of *M. incognita* (Agropolis and Garriga) and
141 three populations of *M. javanica* (Bay, MJ05 and Tugues). Each plant-RKN population
142 combination was repeated 7 and 8 times in the first and second experiment, respectively.
143 The experimental procedures and assessments were those described previously. The
144 experiment was carried out twice

145 *Response of C. metuliferus BGV11135 to virulent RKN populations*

146 The response of the *C. metuliferus* accession BGV11135 and the susceptible melon cv.
147 Paloma was assessed against three *Mil.2* virulent RKN populations belonging to *M.*
148 *arenaria* (MAA106), *M. incognita* (MIA115) and *M. javanica* (MJ27) in 200 cm^{-3} -pot

149 experiments. The avirulent *M. javanica* population MJ05 was included as standard for
150 comparison. The experiment was repeated once. Each plant-RKN population
151 combination was repeated 8 times each experiment. The experimental procedures and
152 assessments were those described previously.

153 *Histopathology*

154 Seeds of *C. metuliferus* BGV11135 and cucumber cv. Dasher II were surface sterilized
155 and incubated as previous to be transferred to transparent envelopes with sterilized paper
156 to maintain humidity for suitable root growth and incubated at 25 ± 2 °C and 16:8 h
157 (light:darkness) photoperiod. Plantlets were inoculated at two true leaf expanded stage
158 with 2500 J2 of *M. javanica* MJ05. After 12 days, roots were carefully washed and cut in
159 pieces of 10 mm. Then, roots containing galls were selected and fixed in 2.5% (v/v)
160 glutaraldehyde in 0.1 M sodium phosphate buffer (pH 7.2) overnight at 4°C and washed
161 three times with same buffer. Afterwards, root pieces were post-fixed in 1% (w/v)
162 osmium tetroxide in 0.1 M sodium phosphate buffer (pH 7.2) for 1 h and washed three
163 times with the same buffer and dehydrated in an acetonitrile series (30–100%) before
164 embedding in epoxy resin (Embed 812, Anamed®) and polymerizing at 60°C for 48h.
165 Semithin (2µm) sections of samples were obtained in a Reichert-Jung Ultracut E Ultra
166 Microtome Leica EM UC6 (Leica Microsysteme GmbH Wien, Austria) and left to dry on
167 a slide previous to be stained with Richardson's blue. The sections were mounted in DPX
168 mountant for histology and observed under a Leica DM4000 B microscope (Leica
169 Microsystems, Mannheim, Germany). Afterwards, sections were photographed using a
170 Leica DFC300 FX 1.4-megapixel digital colour camera equipped with the Leica software
171 application suite LAS V3.8 (Leica Microsystems).

172 *Compatibility and fruit quality assessment*

173 The performance of *C. metuliferus* BGV11135 as rootstock was evaluated using the cv.
174 Vedrantaïs (COMAV) and Paloma (Fitó) of Charentais melon (*Cucumis melo* L. var.
175 *cantalupensis* Naudin) and cv. Finura (Rijk Zwaan) of Piel de sapo melon (*Cucumis melo*
176 L. var. *inodorus* Naudin) as scions. Plants were selfgrafted and grafted onto *C. metuliferus*
177 BGV11135 using the cleft procedure (Lee *et al.*, 2010). Plants were grown under
178 hydroponic conditions in a commercial greenhouse at Fundació Cajamar (Païporta,
179 València) during the spring-summer of 2017. In order to evaluate the impact of grafting
180 on fruit quality, each fruit (8 per treatment) was characterized for the following traits:
181 weight, length and width, rind and flesh thickness, rind and flesh firmness (measured with
182 a digital Penetrometer (8 mm) FHT-803®, Melrose, MA), pH (measured with pH-
183 indicator paper pH1-14 Merck, Darmstadt, Germany), total soluble solids (quantified
184 through a digital Refractometer Atago®, Tokyo, Japan), and flesh color (measured with
185 a colorimeter, Minolta CR-400, New Jersey, USA using the color parameters Hunter L,
186 a and b).

187 *Statistical analysis*

188 Analysis of variance was performed using SAS system V9 (SAS Institute, Inc., Cary, NC,
189 USA). Data on number of eggs masses and eggs per plant were transformed to log₁₀
190 (x+1) when needed to normalize them. The repetitions of the same experiment were
191 compared by the proc glm procedure and consider as the same experiment if no
192 differences were found. Means were separated by the least significant differences (LSD)
193 test when statistical analysis was significant ($P < 0.05$). Paired comparisons between each
194 grafted and selgrafted cultivars for fruit quality traits were done by Student t-test.

195

196 RESULTS

197 *Response of C. metuliferus accessions against avirulent populations of Meloidogyne spp*
198 Both *C. metuliferus* accessions (BGV11135 and BGV10762) responded as highly
199 resistant ($RI < 1\%$) or resistant ($1\% \leq RI \leq 10\%$) to RKN depending on the nematode
200 population. The number of egg masses and eggs per plant material on both *C. metuliferus*
201 accessions were significantly less ($P < 0.05$) than on the cucumber cv. Dasher II
202 irrespective of the *Meloidogyne* specie (Table 2) or the populations assessed (Table 3).

203 The infective and reproductive capacity of *Meloidogyne* populations differed (P
204 < 0.05) on both *C. metuliferus* BGV11135 and cucumber cv. Dasher II (Table 3). The
205 nematode populations Agropolis and Garriga of *M. incognita*, and MJ05 of *M. javanica*
206 produced the highest number of egg masses on *C. metuliferus*, whilst *M. arenaria*
207 population MA68 did on cucumber cv. Dasher II. Regarding RKN reproduction, the *M.*
208 *incognita* population Garriga produced more eggs ($P < 0.05$) than the remaining RKN
209 populations on *C. metuliferus* whilst populations Agropolis and Garriga did on cucumber
210 cv. Dasher II. The accession BGV11135 of *C. metuliferus* performed as resistant against
211 the most RKN populations assessed.

212 *Response of C. metuliferus against virulent Mi1.2 populations of Meloidogyne spp*

213 *C. metuliferus* accession BGV11135 responded as highly resistant ($RI < 1\%$), resistant
214 ($1\% \leq RI \leq 10\%$) or moderately resistant ($10\% \leq RI < 25\%$) to RKN, depending on the
215 nematode population assessed irrespective of its *Mi1.2* gene (a)virulent condition (Table
216 4).

217 *Compatibility and fruit quality assessment*

218 *C. metuliferus* used as rootstock did not affect plant growth of Charentais and Piel de
219 Sapo melons. Grafted plants of each cultivars showed similar vine vigour and flowering
220 time than their respective selfgrafted plants. There was no effect of the rootstock on fruit

221 external and internal quality in the two Charentais melons cultivars, except from a slight
222 increase of flesh thickness in cv. Paloma (Table 5). Each grafted Charentais cultivar
223 maintained fruit size, rind and flesh firmness, and flesh quality (° Brix, pH and colour).
224 Grafting the Piel de sapo melon cv. Finura onto *C. metuliferus* increased the fruit weight
225 and length, but were softer, sweeter and the flesh with lighter colour respect selfgrafted
226 plants (Table 5).

227 *Histopathology*

228 *M. javanica* population MJ05 induced giant cells in both *Cucumis* species (Figure 1) but
229 those produced in *C. metuliferus* were mostly poor developed with multiple vacuoles
230 compared to those on cucumber. Furthermore, giant cells without cytoplasm and necrotic
231 areas surrounding the nematode were observed.

232 DISCUSSION

233 The *C. metuliferus* accessions assessed in this study performed as highly resistant (RI <
234 1%) or resistant (RI = 1% - 10%) to the most RKN populations. These results are in
235 agreement with those reported previously (Fassuliotis, 1967 & 1970; Sigüenza *et al.*,
236 2005; Walters *et al.*, 2006; Thies *et al.*, 2014; De You-Ye *et al.*, 2017). The host suitability
237 of *C. metuliferus* was not affected by the (a)virulent condition of the nematode population.
238 Then, it could be a useful tool to manage RKN nematodes and to prevent the selection of
239 virulent populations in cropping sequences with resistant tomato cultivars or rootstocks.
240 In addition, the *C. metuliferus* accessions assessed in this study are highly resistant to
241 fusarium wilt (Gisbert *et al.*, 2014), and tolerant to *Monosporascus cannonballus* in field
242 conditions (Perpiñà *et al.*, com pers).

243 Fassulotis reported the resistance response of *C. metuliferus* accession C-701 to
244 *M. incognita* in 1967 and conducted histopathological studies in 1970, who observed

245 small giant cells affecting nematode development and increasing the proportion of males.
246 However, no hypersensitive response was observed. Similar results were found by
247 Walters *et al.*, (2006) in the accession PI482454 inoculated with *M. arenaria*, *M. hapla*,
248 *M. incognita* or *M. javanica*. Recently, Ye *et al.*, (2017) have reported a reduction of the
249 number of J2 of *M. incognita* in roots of *C. metuliferus* accession PI482443 at 7 than at 4
250 days after inoculation (dpi), indicating death or emigration from roots and a delayed
251 development of those remaining in them. Empty or poor developed giant cells with
252 multiple vacuoles were observed at 7 and 14 dpi, giant cells appeared to be collapsed or
253 without cytoplasm. In addition, several genes related to plant defence mechanisms were
254 significantly modified and, in contrast with previous reports, hypersensitive necrosis was
255 observed. The results of this study are consistent with those previously reported, in which
256 giant cells produced were multivacuolated and some of them surrounding the nematode
257 area appeared collapsed without cytoplasm. Furthermore, necrotic areas were observed.
258 These results could indicate that the *C. metuliferus* genetic background could play an
259 important role in the interaction with *Meloidogyne sp.*

260 Grafting can affect fruit quality depending on the rootstock-scion interactions,
261 climatic and agronomic conditions (Leonardi *et al.*, 2017). For instance, fruit melons of
262 cv. Supermarket or cv. Proteo grafted onto *C. metuliferus* contained less ° Brix than the
263 ungrafted plants in one out two cropping seasons (Trionfetti-Nisini *et al.*, 2002). Guan *et al.*,
264 (2014) reported less ° Brix content and flesh firmness in galia melons but not from
265 honeydew melons grafted onto *C. metuliferus* conducted in a conventional manner, but
266 not under organic farming. In this study, no differences were found on growth or fruit
267 quality from selfgrafted cantaloupe melon cv. Vedrantaïs and cv. Paloma with those
268 grafted onto *C. metuliferus*. These results are in agreement with those reported by Gisbert
269 *et al.* (2017) who did not find differences among fruit quality from ungrafted, selfgrafted

270 or grafted cv. Vedrantaís onto *C. metuliferus*. Conversely, grafted melon Piel de sapo cv.
271 Finura onto *C. metuliferus* affected fruit weight and length. Nonetheless, these changes
272 do not reduce the commercial value of the fruits as the market of Piel de sapo melons
273 accept a wide range of melon sizes and variability in shapes. The changes in parameters
274 associated with flesh quality (higher ° Brix, lower flesh firmness and lighter flesh color)
275 might be associated to a more advanced ripening state of the grafted melons onto *C.*
276 *metuliferus*. Effects on fruit quality in grafted plants due to growing cycle alterations have
277 been reported previously (Davis *et al.*, 2008; Soteriou *et al.*, 2014). Therefore, these
278 effects could be reduced adapting the harvesting period for each rootstock-scion
279 combination.

280 In conclusion, *C. metuliferus* accession BGV11135 could be a promising melon
281 rootstock to manage *Meloidogyne spp.* irrespective of its (a)virulent *Mi1.2* condition
282 without melon fruit quality reduction.

283

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288

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